

Please amend Claims 1, 3, 5 and 10, as follows:

B1
1. (Amended) A nucleotide sequence comprising, a regulatory region in operative association with a xylanase secretion sequence and a gene of interest, wherein [at least one of the regulatory region, or] the gene of interest is not [normally] associated with the production of a xylanase protein that is native to said xylanase secretion signal.

B2
3. (Amended) The nucleotide sequence of claim 1, wherein the gene of interest is selected from a gene encoding a protein selected from the group consisting of a pharmaceutical, a nutraceutical, an industrial enzyme, an animal feed, a food additive, and an enzyme.

B3
5. (Amended) The nucleotide sequence of claim 1 further comprising a selectable marker [gene].

B4
10. (Amended) The transformed filamentous fungi of claim 9, wherein the filamentous fungi is selected from the group consisting of *Trichoderma*, *Humicola*, *Fusarium*, *Aspergillus*, *Mycogone*, *Verticillium*, [*Streptomyces*], *Colletotrichum*, *Neurospora*, *Botrytis*, *Pleurotus*, *Penicillum*, *Cephalosporium*, *Myrothecium*, *Papulospora*, *Achlya*, *Podospora*, *Endothia*, *Mucor*, *Cochilobolus*, *Tolypocladium*, *Pyricularia*, *Penicillium*, *Myceliophthora*, *Irpex*, *Stachybotrys*, *Scorulariopsis*, *Chaetomium*, [*Gilocladium*], *Gliocladium*, *Cephalosporin* and *Acremonium*.

Please add new Claims 31-33, as follows: